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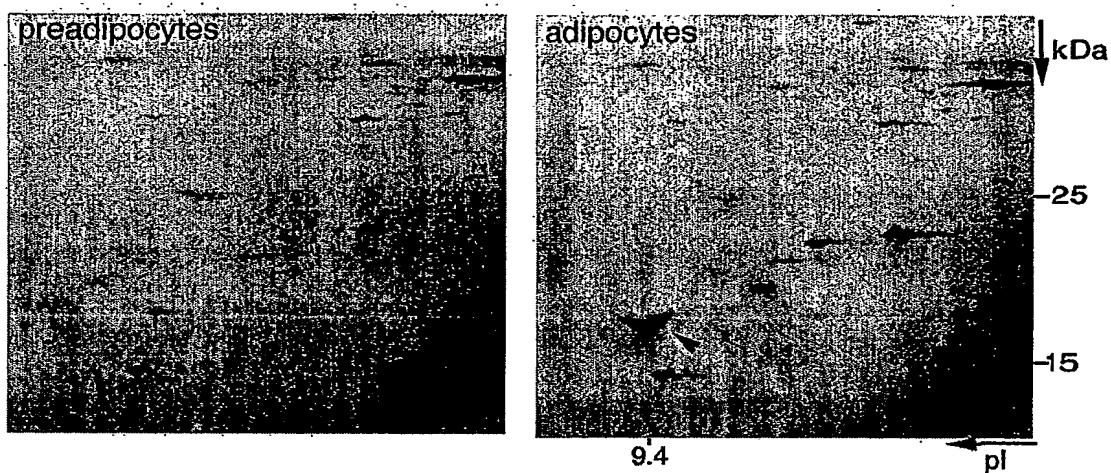
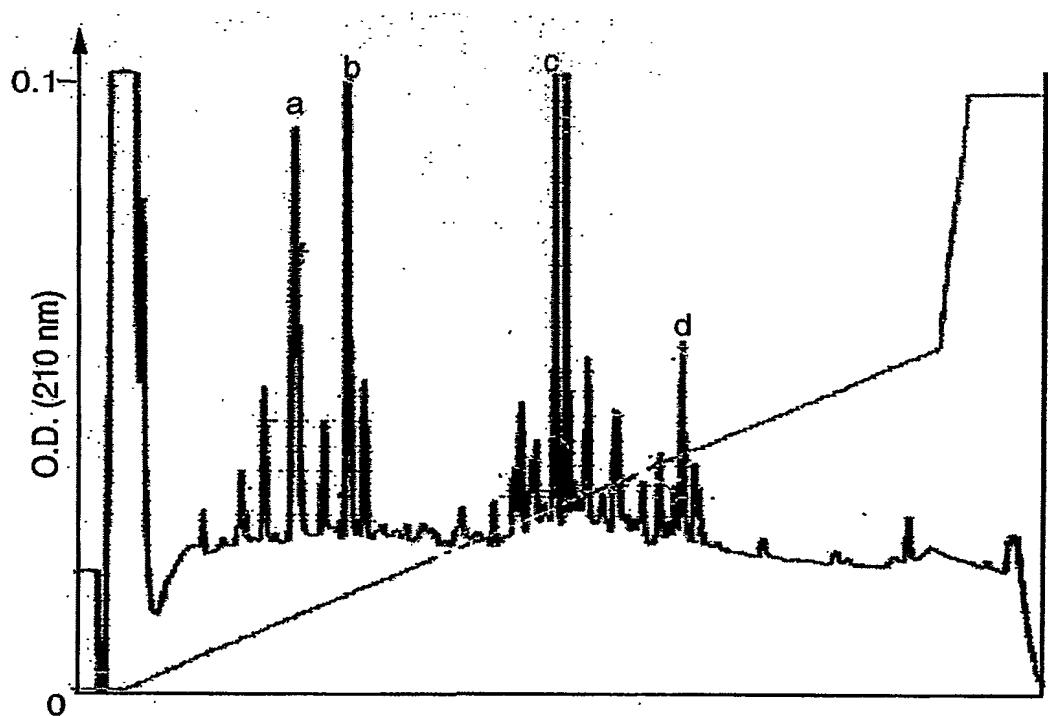


Figure 1

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Fractions	Amino acid sequences
a	LQQTNCPK
b	GTEPELSETQR
c	CLLISLALWLGTVGTR
d	HPPVQLAFQEIGVDR

Figure 2

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A. [REDACTED]

MKGCLMISIATLWGGTVGIRGTEPELSETQRRSLQVALEEFHKHPPVQLAFQEIGVDRAEEVLFSAGTFVRLEFKLQQTNCPKKDWKKPECTIKPNGRRKCLACIKMDPKGKILGRIVHCPILKQGPQDPQELQCIKIAQAGEDPHGYFLPGQFAFSRALRTK

B.

SS (1-17)	Potential cystatin like domain (32-108)	Domain with unknown function (109-162)
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C.

Adipocyspin

LQV[LE][FH]HPPVQLAFQEIGVDR[EEVLFS][FVR][EFKLQQ][N][P][KDWKKP]-IT
90

Cystatin C

ADFAVA[YN][KSNDLYELRV][K][VS][KSQVV-[A-C][NYY][KVEVGE][E][K][LSPTDL][N][P] 78

Query: 91

IKPN[RRRK][LACIKMD] 108

Sbjct: 79

FLEQ[EEGF][DFEVWEK] 96

Figure 3